| Project Title | Funding | Strategic Plan Objective | Institution | |
|---|-------------|--------------------------|--|--|
| Advanced parental age and autism: The role of aneuploidy and uniparental disomy in ASD pathogenesis | \$0 | Q3.S.A | Albert Einstein College of Medicine of Yeshiva Universit | |
| Autism Genome Project (AGP) | \$10,000 | Q3.L.B | Autism Speaks (AS) | |
| A genome-wide search for autism genes in the SSC Baylor | \$20,344 | Q3.L.B | Baylor College of Medicine | |
| Simons Simplex Collection Site | \$0 | Q3.L.B | Baylor College of Medicine | |
| Simons Simplex Collection support grant | \$30,000 | Q3.L.B | Baylor College of Medicine | |
| Finding recessive genes for autism spectrum disorders | \$349,999 | Q3.L.B | Boston Children's Hospital | |
| Simons Simplex Collection Site | \$51,656 | Q3.L.B | Boston Children's Hospital | |
| Simons Simplex Collection support grant | \$30,000 | Q3.L.B | Boston Children's Hospital | |
| A genome-wide search for autism genes in the SSC CHB | \$50,000 | Q3.L.B | Boston Children's Hospital | |
| Autism genetics: Homozygosity mapping and functional validation | \$850,815 | Q3.S.A | Boston Children's Hospital | |
| A genome-wide search for autism genes in the SSC Brown | \$50,000 | Q3.L.B | Brown University | |
| Understanding the genetic basis of autism | \$6,557,422 | Q3.L.B | Cold Spring Harbor Laboratory | |
| Genetic basis of autism | \$0 | Q3.L.B | Cold Spring Harbor Laboratory | |
| Novel statistical methods for DNA sequencing data, and applications to autism | \$339,743 | Q3.L.B | Columbia University | |
| Simons Simplex Collection Site | \$0 | Q3.L.B | Columbia University | |
| Simons Simplex Collection support grant | \$1,430 | Q3.L.B | Columbia University | |
| Simons Simplex Collection Site | \$0 | Q3.L.B | Emory University | |
| A genome-wide search for autism genes in the SSC Emory | \$72,524 | Q3.L.B | Emory University | |
| Simons Simplex Collection support grant | \$30,682 | Q3.L.B | Emory University | |
| Population genetics to improve homozygosity mapping and mapping in admixed groups | \$52,190 | Q3.L.B | Harvard Medical School | |
| Illumina, Inc. | \$717,504 | Q3.L.B | Illumina, Inc. | |
| Genomic profiling of autism families using whole- genome sequencing | \$129,600 | Q3.L.B | Institut Pasteur | |
| Integrative genetic analysis of autistic brains | \$200,000 | Q3.L.B | Johns Hopkins University School of Medicine | |
| Cryptic chromosomal aberrations contributing to autism | \$70,524 | Q3.L.B | Massachusetts General Hospital | |
| Complex genetic architecture of chromosomal aberrations in autism | \$92,917 | Q3.L.B | Massachusetts General Hospital | |
| Rapid characterization of balanced genomic rearrangements contributing to autism | \$53,942 | Q3.L.B | Massachusetts General Hospital | |
| Sequence-based discovery of genes with pleiotropic effects across diagnostic boundaries and throughout the lifespan | \$0 | Q3.L.B | Massachusetts General Hospital and Harvard University | |

| Project Title | Funding | Strategic Plan Objective | Institution |
|--|-------------|--------------------------|---|
| Simons Simplex Collection support grant | \$30,040 | Q3.L.B | McGill University Health Centre- Montreal Children's Hospital |
| Hypocholesterolemic autism spectrum disorder | \$84,549 | Q3.L.B | National Institutes of Health |
| Genetic epidemiology of complex traits | \$559,192 | Q3.L.B | National Institutes of Health |
| Autism Genome Project (AGP) Core Consortium | \$0 | Q3.L.B | Nationwide Children's Hospital |
| Mitochondria and the etiology of autism | \$437,500 | Q3.L.B | The Children's Hospital of Philadelphia |
| Autism Genome Project (AGP): Genome sequencing and analysis supplement | \$50,000 | Q3.L.B | The Hospital for Sick Children |
| Examining the Y-chromosome in autism spectrum disorder | \$175,000 | Q3.L.B | The Hospital for Sick Children |
| Identifying genetic variants on the Y chromosome of males with autism | \$50,555 | Q3.L.B | The Hospital for Sick Children |
| The frequency of polymorphisms in maternal- and paternal-effect genes in autism spectrum | \$152,545 | Q3.L.B | The Pennsylvania State University |
| Simons Simplex Collection Site | \$44,598 | Q3.L.B | The Research Institute of the McGill University Health Centre |
| Autism Genome Project Consortium data reanalysis using computational biostatistics | \$60,000 | Q3.L.B | The Rockefeller University |
| Genomic influences on development and outcomes in infants at risk for autism | \$498,341 | Q3.L.B | University of Alberta |
| Genomic influences on development and outcomes in Infants at risk of ASD | \$0 | Q3.S.A | University of Alberta |
| Genomic influences on developmental course and outcome in Infants at risk of ASD: A Baby Siblings Research Consortium (BSRC) Study | \$147,661 | Q3.S.A | University of Alberta |
| Simons Simplex Collection Site | \$0 | Q3.L.B | University of California, Los Angeles |
| A genome-wide search for autism genes in the SSC UCLA | \$100,000 | Q3.L.B | University of California, Los Angeles |
| Simons Simplex Collection support grant | \$30,000 | Q3.L.B | University of California, Los Angeles |
| ACE Network: A comprehensive approach to identification of autism susceptibility genes (supplement) | \$442,627 | Q3.L.B | University of California, Los Angeles |
| ACE Network: A comprehensive approach to identification of autism susceptibility genes | \$2,631,440 | Q3.L.B | University of California, Los Angeles |
| Genome-wide expression profiling data analysis to study autism genetic models | \$0 | Q3.S.A | University of California, Los Angeles |
| Rapid phenotyping for rare variant discovery in autism | \$700,956 | Q3.S.A | University of California, Los Angeles |
| Whole-exome sequencing to identify causative genes for autism | \$350,000 | Q3.L.B | University of California, San Diego |
| Dissecting expression regulation of an autism GWAS hit | \$30,000 | Q3.L.B | University of California, San Francisco |
| Linking autism and congenital cerebellar malformations | \$0 | Q3.L.B | University of Chicago |

| Project Title | Funding | Strategic Plan Objective | Institution | |
|---|-------------|--------------------------|---|--|
| Investigation of DUF1220 domains in human brain function and disease | \$376,668 | Q3.L.B | University of Colorado Denver | |
| Simons Simplex Collection support grant | \$30,000 | Q3.L.B | University of Illinois at Chicago | |
| Simons Simplex Collection Site | \$0 | Q3.L.B | University of Illinois at Chicago | |
| A genome-wide search for autism genes in the SSC UIC | \$48,419 | Q3.L.B | University of Illinois at Chicago | |
| Simons Simplex Collection Site | \$123,678 | Q3.L.B | University of Michigan | |
| Simons Simplex Collection Site | \$0 | Q3.L.B | University of Missouri | |
| Simons Simplex Collection support grant | \$30,000 | Q3.L.B | University of Missouri | |
| 3/3-Sequencing autism spectrum disorder extended pedigrees | \$160,000 | Q3.L.B | University of Pennsylvania | |
| A history of behavioral genetics | \$0 | Q3.Other | University of Pittsburgh | |
| A genome-wide search for autism genes in the SSC Pittsburgh | \$50,000 | Q3.L.B | University of Pittsburgh | |
| Autism Genome Project (AGP) Core Consortium | \$0 | Q3.L.B | University of Pittsburgh | |
| FOXP2-regulated signaling pathways critical for higher cognitive functions | \$248,921 | Q3.Other | University of Texas Southwestern Medical Center | |
| FOXP2-regulated signaling pathways critical for higher cognitive functions (supplement) | \$66,686 | Q3.Other | University of Texas Southwestern Medical Center | |
| 1/3-Sequencing autism spectrum disorder extended pedigrees | \$299,000 | Q3.L.B | University of Utah | |
| Simons Simplex Collection support grant | \$29,752 | Q3.L.B | University of Washington | |
| Simons Simplex Collection Site | \$75,000 | Q3.L.B | University of Washington | |
| Genomic hotspots of autism | \$261,033 | Q3.L.B | University of Washington | |
| Whole exome sequencing of Simons Simplex Collection quads | \$1,835,440 | Q3.L.B | University of Washington | |
| 2/3-Sequencing autism spectrum disorder extended pedigrees | \$231,688 | Q3.L.B | University of Washington | |
| Next generation gene discovery in familial autism | \$688,392 | Q3.L.B | University of Washington | |
| Simons Simplex Collection Site | \$0 | Q3.L.B | Vanderbilt University | |
| Simons Simplex Collection support grant | \$30,000 | Q3.L.B | Vanderbilt University Medical Center | |
| A genome-wide search for autism genes in the SSC Vanderbilt | \$300,000 | Q3.L.B | Vanderbilt University Medical Center | |
| Simons Foundation Simplex Project Collection Site | \$0 | Q3.L.B | Weill Cornell Medical College | |
| Simons Simplex Collection support grant | \$34,200 | Q3.L.B | Weill Cornell Medical College | |
| A genome-wide search for autism genes in the Simons Simplex Collection | \$415,782 | Q3.L.B | Yale University | |
| Whole exome sequencing of Simons Simplex Collection quads | \$2,110,073 | Q3.L.B | Yale University | |

| Project Title | Funding | Strategic Plan Objective | Institution |
|--|-----------|--------------------------|-----------------|
| Whole Exome Sequencing of Simons Simplex Trios | \$114,106 | Q3.L.B | Yale University |
| ACE Center: Rare variant genetics, contactin-related proteins and autism | \$324,189 | Q3.L.B | Yale University |
| Simons Simplex Collection Site | \$96,641 | Q3.L.B | Yale University |
| Simons Simplex Collection support grant | \$30,000 | Q3.L.B | Yale University |